## AMENDMENTS TO THE CLAIMS:

This listing of the claims will replace all prior versions, and listings, of claims in the present application.

## LISTING OF THE CLAIMS:

- 31. (currently amended) A method of isolating at least one anti-ligand to at least one target ligand comprising the steps of:
  - (i) providing a library of anti-ligands;
- (ii) providing an amount of a first subtractor ligand a first population of ligands comprising a ligand fixed to or incorporated in a subtractor ligand construct;
- (iii) providing an amount of a second target ligand a second population of ligands comprising the same ligand as step (ii), fixed to or incorporated in a target ligand construct;
  - (iv) determining amounts of first subtractor <u>ligand construct</u> and seeond target <del>ligand construct</del> using one or more equations derived from the universal law of mass

action 
$$\frac{[C]^c[d]^d}{[A]^a[B]^b} = K_{eq},$$

where

A, B, C & D = are the participants in the reaction (reactants and products)

a, b, c, & d = the coefficients necessary for a balanced chemical equation

- so as to permit isolation of at least one anti-ligand to at least one target ligand;

  (v) providing the amount of the first subtractor ligand construct determined in step (iv);
- (vi) providing the amount of the second target ligand construct determined in step (iv):
- (vii) providing separation means for isolating anti-ligand bound to the seeend target ligand <u>construct</u> from anti-ligand bound to the <u>first</u> subtractor ligand <u>construct</u>, <u>wherein said</u> <u>separation means for the subtractor and target ligand constructs can be either the same or</u> different;
- (viii) exposing the library of (i) to the ligands of (v) and (vi) to permit binding of antiligands to ligands; and
- (ix) isolating the anti-ligand bound to the second ligand fixed to or incorporated in the target ligand construct with using the separation means.

Claim 32-33 (canceled)

- 34. (currently amended) A method as claimed in one-of-claims claim 31, 32-or-33 comprising a further step of releasing the anti-ligand from the second-target ligand.
- 35. (currently amended) A method as claimed in claim 31 er 32 whereby steps (ii) to (ix) are repeated one or more times.
- 36. (currently amended) A method as claimed in claim 31 or 32 wherein the amount of one of the first subtractor ligand construct or second target ligand construct is provided in excess of the amount of the other of the first subtractor ligand construct or second the target ligand construct.
- 37. (previously presented) A method as claimed in claim 36 where the excess of ligand is of 10 to 100 fold.
- 38. (currently amended) A method as claimed in elaims claim 31 or 32 wherein the equation of (iv) is

$$bA = \frac{\left(A + T + (K_d)x(CxV)\right)}{2} - \sqrt{\frac{\left(A + T + (K_d)x(CxV)\right)^2}{4} - AxT}$$

where

bA = Bound anti-ligand

A = Total number of anti-ligand

T = Total number of ligands

C = Avogadro's constant (6.022 x 10<sup>23</sup> particles/mole)

V = Reaction volume (litres)

 $K_d$  = Equilibrium dissociation constant.

39. (currently amended) A method as claimed in claims 31 or 32 wherein the equation of (iv) is:

Application No. 10/526,695 February 18, 2009

$$b\mathcal{A} = \left\{ \frac{\left(A + T + (K_g)x(CxV)\right)}{2} - \sqrt{\frac{\left(A + T + (K_g)x(CxV)\right)^2}{4} \cdot AxT} \right\} x \left\{ \frac{\left(T_gxC_g\right)}{\left(\left(T_gxC_g\right) + \left(T_gxC_g\right)\right)} \right\}$$

where

bAn = Bound anti-ligand

Tp = The number of ligands on Cp

 $T_s =$ The number of ligands on  $C_s$ 

Cp = The number of target ligand constructs

Cs = The number of subtractor ligand constructs

A = Total number of anti-ligand

T = Total number of ligands

C = Avogadro's constant (6.022 x 10<sup>23</sup> particles/mole)

V = Reaction volume (litres)

K<sub>d</sub> = Equilibrium dissociation constant.

- 40. (currently amended) A method as claimed in claim 31 e<del>x 32</del> wherein the separation means are selected from at least one of a solid support, cell membrane and/or portions thereof, synthetic membrane, beads, chemical tags and free ligand.
- 41.(previously presented) A method as claimed in claim 40 whereby the separation means are cell membranes and/or portions thereof.
- 42. (previously presented) A method as claimed in claim 41 whereby the first subtractor and seeond target ligands are fixed to and/or incorporated within separate cell membranes and/or portions thereof.
- 43. (currently amended) A method as claimed in elaims <u>claims</u> 31 or 32 whereby the separation means of the first-subtractor and seeond target ligand <u>constructs</u> have a different density.
  - 44. (previously presented) A method as claimed in claim 43 wherein the separation

Application No. 10/526,695 February 18, 2009

means of the first subtractor ligand <u>construct</u> is of a lower density than the separation means of the second target ligand construct.

- 45.(previously presented) A method as claimed in claim 44 wherein the separation means of the first subtractor ligand construct is a membrane vesicle.
- 46. (previously presented) A method as claimed in claim 44 wherein the separation means of the second target ligand <u>construct</u> is a whole cell membrane.
- 47. (currently amended) A method as claimed in claim 31 or 32 whereby the isolation of anti-ligand bound to second target ligand step (ix) is performed by at least one of density centrifugation, solid support sequestration, magnetic bead sequestration, chemical tag binding and aqueous phase partitioning.
- 48. (previously presented) A method as claimed in claim 47 whereby the isolation step is performed by density centrifugation.
- 49. (previously presented) A method as claimed in claim 48 wherein the density centrifugation is performed using a sucrose-polymer gradient.
- 50. (currently amended) A method as claimed in claim 31 or 32 wherein the library of step (i) is a display library comprising a plurality of library members which display anti-ligands.
- 51. (previously presented) A method as claimed in claim 50 wherein the library is a phage display library.
- 52. (currently amended) A method as claimed in claim 31 er-32 wherein the subtractor and target ligands are independently at least one from antigens; receptor ligands; and enzyme targets that comprise at least one selected from carbohydrate; protein; peptide; lipid; polynucleotide; inorganic molecules and conjugated molecules.

Application No. 10/526,695 12578/46501 February 18, 2009

53. (currently amended) A method as claimed in claim 31 or 32-wherein the library of anti-ligands is composed of at least one selected from antibodies, and antigen binding variants, derivatives or fragments thereof; scaffold molecules with engineered variable surfaces; receptors; and enzymes.

54. (currently amended) A method as claimed in claim 31 or 32 comprising a further step of exposing the ligand and its separation means to a stimulus which influences the expression of target ligands on said ligand constructs.

Claims 55-59 (canceled)